

	Type	L #	Hits	Search Text
2	BRS	L2	71	(dementia and acetylcholinesterase\$.clm.

*Interference Search*  
*10/22/07*  
*[Signature]*

=> d his

(FILE 'HOME' ENTERED AT 09:27:09 ON 22 OCT 2007)

FILE 'REGISTRY' ENTERED AT 09:27:25 ON 22 OCT 2007

L1 SCREEN 2039 AND 1842 AND 2004 AND 1994 AND 1949  
 L2 STRUCTURE UPLOADED  
 L3 QUE L2 AND L1  
 L4 0 S L3  
 L5 0 S L3 FULL  
 L6 SCREEN 2039 AND 1842 AND 2004 AND 1994 AND 1949  
 L7 STRUCTURE UPLOADED  
 L8 QUE L7 AND L6  
 L9 0 S L8  
 L10 0 S L8 FULL  
 L11 SCREEN 2039 AND 1842 AND 2004 AND 1994 AND 1949  
 L12 STRUCTURE UPLOADED  
 L13 QUE L12 AND L11  
 L14 0 S L13  
 L15 1 S L13 FULL

FILE 'CAPLUS' ENTERED AT 09:30:18 ON 22 OCT 2007

L16 3 S L15  
 L17 3 DUP REM L16 (0 DUPLICATES REMOVED)

FILE 'STNGUIDE' ENTERED AT 09:31:23 ON 22 OCT 2007

FILE 'CAPLUS' ENTERED AT 09:32:06 ON 22 OCT 2007

=> d stat que

L11 SCR 2039 AND 1842 AND 2004 AND 1994 AND 1949  
 L12 STR

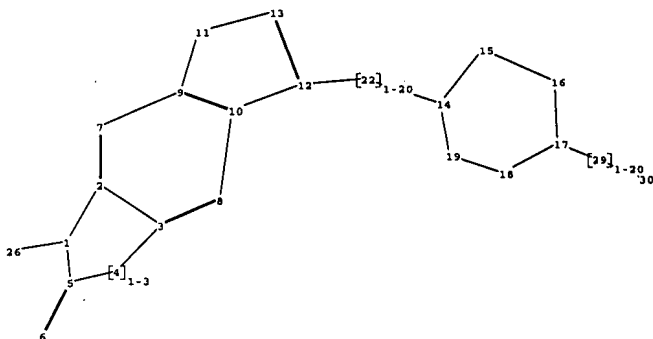
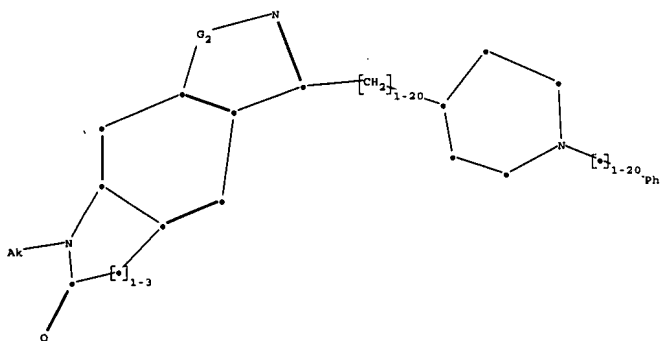
\* STRUCTURE DIAGRAM TOO LARGE FOR DISPLAY - AVAILABLE VIA OFFLINE PRINT \*

Structure attributes must be viewed using STN Express query preparation.

L15 1 SEA FILE=REGISTRY SSS FUL L12 AND L11  
 L16 3 SEA FILE=CAPLUS ABB=ON PLU=ON L15  
 L17 3 DUP REM L16 (0 DUPLICATES REMOVED)

=> log y

*10/22/07  
Interference Search*



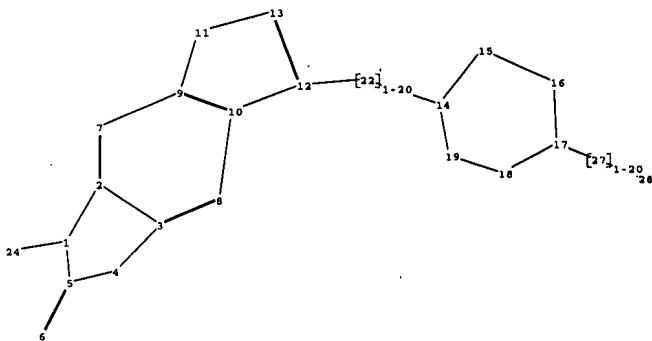
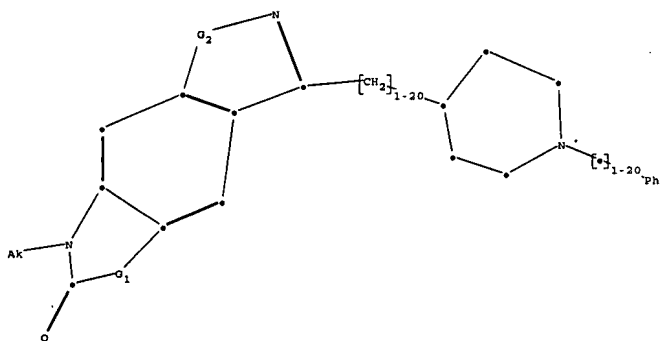
chain nodes :  
6 22 26 29 30  
ring nodes :  
1 2 3 4 5 7 8 9 10 11 12 13 14 15 16 17 18 19  
chain bonds :  
1-26 5-6 12-22 14-22 17-29 29-30  
ring bonds :  
1-2 1-5 2-3 2-7 3-4 3-8 4-5 7-9 8-10 9-10 9-11 10-12 11-13 12-13 14-15 14-19 15-16 16-17  
17-18 18-19  
exact/norm bonds :  
1-2 1-5 1-26 3-4 4-5 5-6 9-11 10-12 11-13 12-13 12-22 14-15 14-19 14-22 15-16 16-17 17-18  
17-29 18-19 29-30  
normalized bonds :  
2-3 2-7 3-8 7-9 8-10 9-10  
isolated ring systems :  
containing 1 :

G1:C,O,S,N

G2:O,S

Match level :  
1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:CLASS7:Atom 8:Atom 9:Atom 10:Atom 11:Atom 12:Atom  
13:Atom

14:Atom 15:Atom 16:Atom 17:Atom 18:Atom 19:Atom 22:CLASS26:CLASS29:CLASS30:CLASS



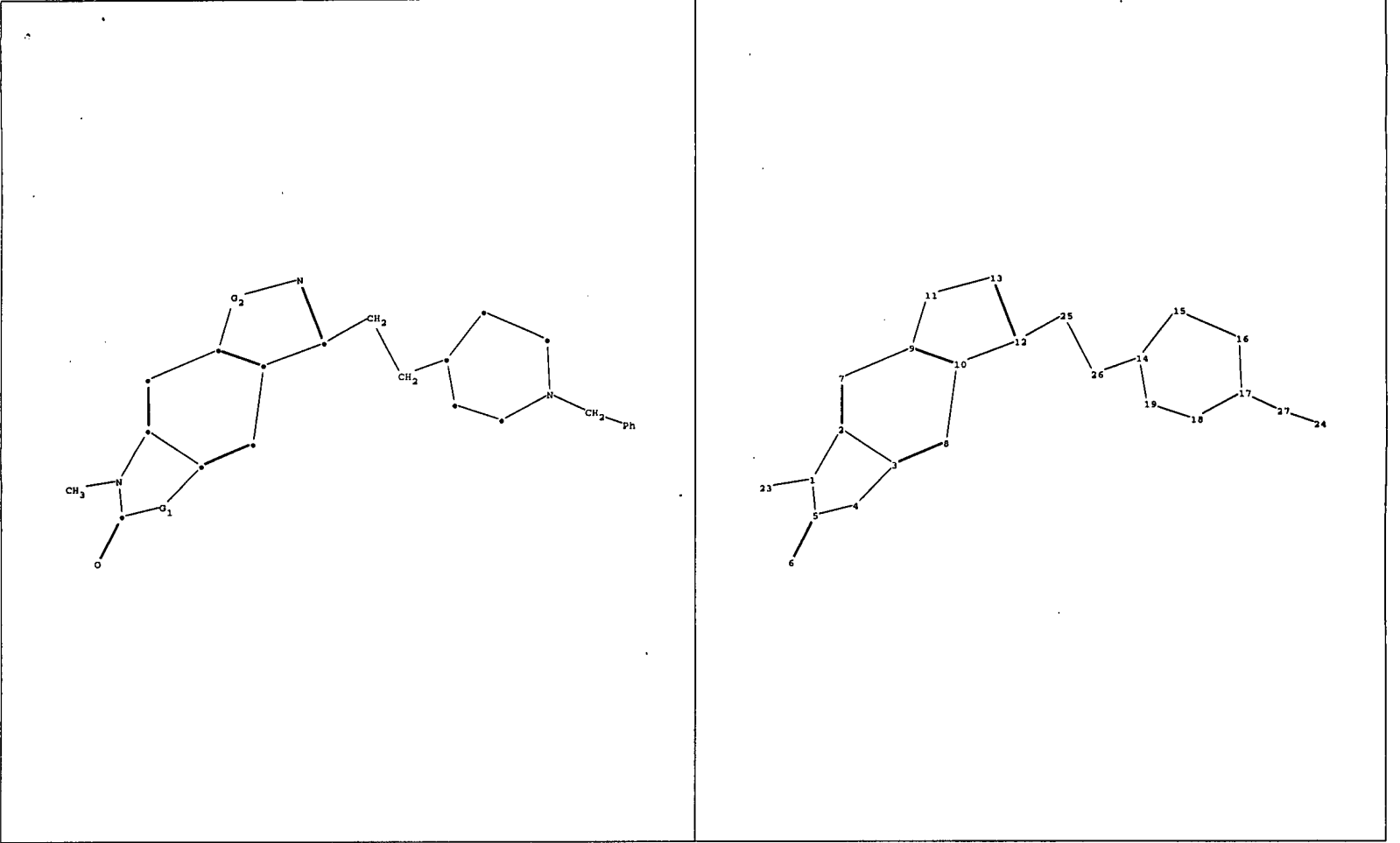
chain nodes :  
6 22 24 27 28  
ring nodes :  
1 2 3 4 5 7 8 9 10 11 12 13 14 15 16 17 18 19  
chain bonds :  
1-24 5-6 12-22 14-22 17-27 27-28  
ring bonds :  
1-2 1-5 2-3 2-7 3-4 3-8 4-5 7-9 8-10 9-10 9-11 10-12 11-13 12-13 14-15 14-19 15-16 16-17  
17-18 18-19  
exact/norm bonds :  
1-2 1-5 1-24 3-4 4-5 5-6 9-11 10-12 11-13 12-13 12-22 14-15 14-19 14-22 15-16 16-17 17-18  
17-27 18-19 27-28  
normalized bonds :  
2-3 2-7 3-8 7-9 8-10 9-10  
isolated ring systems :  
containing 1 :

G1:O,S,N

G2:O,S

Match level :  
1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:CLASS7:Atom 8:Atom 9:Atom 10:Atom 11:Atom 12:Atom  
13:Atom

14:Atom 15:Atom 16:Atom 17:Atom 18:Atom 19:Atom 22:CLASS24:CLASS27:CLASS28:CLASS



chain nodes :  
6 23 24 25 26 27

ring nodes :  
1 2 3 4 5 7 8 9 10 11 12 13 14 15 16 17 18 19

chain bonds :  
1-23 5-6 12-25 14-26 17-27 24-27 25-26

ring bonds :  
1-2 1-5 2-3 2-7 3-4 3-8 4-5 7-9 8-10 9-10 9-11 10-12 11-13 12-13 14-15 14-19 15-16 16-17 17-18 18-19

exact/norm bonds :  
1-2 1-5 1-23 3-4 4-5 5-6 9-11 10-12 11-13 12-13 12-25 14-15 14-19 14-26 15-16 16-17 17-18 17-27 18-19 24-27 25-26

normalized bonds :  
2-3 2-7 3-8 7-9 8-10 9-10

isolated ring systems :  
containing 1 :

G1:O,S,N

G2:O,S

Match level :  
1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:CLASS7:Atom 8:Atom 9:Atom 10:Atom 11:Atom 12:Atom 13:Atom

14:Atom 15:Atom 16:Atom 17:Atom 18:Atom 19:Atom 23:CLASS24:CLASS25:CLASS26:CLASS  
27:CLASS